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SCHUELKE, Norbert
GARDNER, Jason
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<210> 14

<211> 481

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide

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<210> 15

<211> 142

<212> PRT

<213> Homo sapiens

<400> 15

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1 5 10 15

Val Gln Cys Gln Val Gln Leu Val Glu Ser Gly Gly Gly Val Val Gln
20 25 30

Pro Gly Arg Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Ala Phe
35 40 45

Ser Arg Tyr Gly Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu
50 55 60

Glu Trp Val Ala Val Ile Trp Tyr Asp Gly Ser Asn Lys Tyr Tyr Ala
65 70 75 80

Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn
85 90 95

Thr Gln Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val
100 105 110

Tyr Tyr Cys Ala Arg Gly Gly Asp Phe Leu Tyr Tyr Tyr Tyr Tyr Gly
115 120 125

Met Asp Val Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser
130 135 140

<210> 16

<211> 463

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide

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cagagtcacc atcacttgcc gggcgagtca gggcattagc aattatttag cctggtatca 180

gcagaaaaca gggaaagttc ctaagttcct gatctatgaa gcatccactt tgcaatcagg 240

ggtcccatct cggttcagtg gcggtggatc tgggacagat ttcactctca ccatcagcag 300

cctgcagcct gaagatgttg caacttatta ctgtcaaaat tataacagtg ccccattcac 360

tttcggccct gggaccaaag tggatatcaa acgaactgtg gctgcaccct ctgtcttcat 420

cttcccgcca tctgatgagc agttgaaatc tggaactgct agc 463

<210> 17

<211> 127

<212> PRT

<213> Homo sapiens

<400> 17

Met Arg Val Pro Ala Gln Leu Leu Gly Leu Leu Leu Leu Trp Leu Pro
1 5 10 15

Asp Thr Arg Cys Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser
20 25 30

Ala Ser Val Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Gly
35 40 45

Ile Ser Asn Tyr Leu Ala Trp Tyr Gln Gln Lys Thr Gly Lys Val Pro
50 55 60

Lys Phe Leu Ile Tyr Glu Ala Ser Thr Leu Gln Ser Gly Val Pro Ser
65 70 75 80

Arg Phe Ser Gly Gly Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser
85 90 95

Ser Leu Gln Pro Glu Asp Val Ala Thr Tyr Tyr Cys Gln Asn Tyr Asn
100 105 110

Ser Ala Pro Phe Thr Phe Gly Pro Gly Thr Lys Val Asp Ile Lys
115 120 125

<210> 18

<211> 508

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide

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cgtggtccag cctgggaggt ccctgagact ctcctgtgca gcgtctggat tcaccttcag 180

taactatgtc atgcactggg tccgccaggc tccaggcaag gggctggagt gggtggcaat 240

tatatggtat gatggaagta ataaatacta tgcagactcc gtgaagggcc gattcaccat 300

ctccagagac aattccaaga acacgctgta tctgcaaatg aacagcctga gagccgagga 360

cacggctgtg tattactgtg cgggtggata taactggaac tacgagtacc actactacgg 420

tatggacgtc tggggccaag ggaccacggt caccgtctcc tcagcctcca ccaagggccc 480

atcggtcttc cccctggcac cctctagc 508

<210> 19

<211> 143

<212> PRT

<213> Homo sapiens

<400> 19

Met Glu Leu Gly Leu Arg Trp Val Leu Leu Val Ala Leu Leu Arg Gly
1 5 10 15

Val Gln Cys Gln Val Gln Leu Val Glu Ser Gly Gly Gly Val Val Gln
20 25 30

Pro Gly Arg Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe
35 40 45

Ser Asn Tyr Val Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu
50 55 60

Glu Trp Val Ala Ile Ile Trp Tyr Asp Gly Ser Asn Lys Tyr Tyr Ala
65 70 75 80

Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn
85 90 95

Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val
100 105 110

Tyr Tyr Cys Ala Gly Gly Tyr Asn Trp Asn Tyr Glu Tyr His Tyr Tyr
115 120 125

Gly Met Asp Val Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser
130 135 140

<210> 20

<211> 463

<212> DNA

<213> Artificial Sequence
<220>

<223> Synthetic Oligonucleotide

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cagagtcacc atcacttgtc gggcgagtca gggcattacc aattatttag cctggtttca 180

gcagaaacca gggaaagccc ctaagtccct tatctatgct gcatccagtt tgcaaagtgg 240

ggtcccatca aagttcagcg gcagtggatc tgggacagat ttcagtctca ccatcagcag 300

cctgcagcct gaagattttg caacttatta ctgccaacag tataatagtt acccgatcac 360

cttcggccaa gggacacgac tggagattaa acgaactgtg gctgcaccat ctgtcttcat 420

cttcccgcca tctgatgagc agttgaaatc tggaactgct agc 463

<210> 21

<211> 127

<212> PRT

<213> Homo sapiens

<400> 21

Met Arg Val Pro Ala Gln Leu Leu Gly Leu Leu Leu Leu Cys Phe Pro
1 5 10 15

Gly Ala Arg Cys Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser
20 25 30

Ala Ser Val Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Gly
35 40 45

Ile Thr Asn Tyr Leu Ala Trp Phe Gln Gln Lys Pro Gly Lys Ala Pro
50 55 60

Lys Ser Leu Ile Tyr Ala Ala Ser Ser Leu Gln Ser Gly Val Pro Ser
65 70 75 80

Lys Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Ser Leu Thr Ile Ser
85 90 95

Ser Leu Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Tyr Asn
100 105 110

Ser Tyr Pro Ile Thr Phe Gly Gln Gly Thr Arg Leu Glu Ile Lys
115 120 125

<210> 22

<211> 490

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide

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gagactctcc tgtgcagcgt ctggattcac cttcagtagc tatggcatgc actgggtccg 180

ccaggctcca ggcaaggggc tggactgggt ggcaattatt tggcatgatg gaagtaataa 240

atactatgca gactccgtga agggccgatt caccatctcc agagacaatt ccaagaagac 300

gctgtacctg caaatgaaca gtttgagagc cgaggacacg gctgtgtatt actgtgcgag 360

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agggaccacg gtcaccgtct cctcagcctc caccaagggc ccatcggtct tccccctggc 480

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<210> 23

<211> 145

<212> PRT

<213> Homo sapiens

<400> 23

Met Glu Leu Gly Leu Ser Trp Val Phe Leu Val Ala Leu Leu Arg Gly
1 5 10 15

Val Gln Cys Gln Val Gln Leu Val Glu Ser Gly Gly Gly Val Val Gln
20 25 30

Pro Gly Arg Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe
35 40 45

Ser Ser Tyr Gly Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu
50 55 60

Asp Trp Val Ala Ile Ile Trp His Asp Gly Ser Asn Lys Tyr Tyr Ala
65 70 75 80

Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Lys
85 90 95

Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val
100 105 110

Tyr Tyr Cys Ala Arg Ala Trp Ala Tyr Asp Tyr Gly Asp Tyr Glu Tyr
115 120 125

Tyr Phe Gly Met Asp Val Trp Gly Gln Gly Thr Thr Val Thr Val Ser
130 135 140

Ser
145

<210> 24

<211> 463

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide

<400> 24
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tgccagatgt gacatccaga tgacccagtc tccatcctca ctgtctgcat ctgtaggaga 120

cagagtcacc atcacttgtc gggcgagtca gggcattagc cattatttag cctggtttca 180

gcagaaacca gggaaagccc ctaagtccct gatctatgct gcatccagtt tgcaaagtgg 240

ggtcccatca aagttcagcg gcagtggatc tgggacagat ttcactctca ccatcagcag 300

cctacagcct gaagattttg caacttatta ctgccaacag tataatagtt tcccgctcac 360

tttcggcgga gggaccaagg tggagatcaa acgaactgtg gctgcaccat ctgtcttcat 420

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<210> 25

<211> 127

<212> PRT

<213> Homo sapiens

<400> 25

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Met Arg Val Pro Ala Gln Leu Leu Gly Leu Leu Leu Leu Cys Phe Pro
1               5                  10                  15

Gly Ala Arg Cys Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser
            20                  25                  30

Ala Ser Val Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Gly
        35                  40                  45

Ile Ser His Tyr Leu Ala Trp Phe Gln Gln Lys Pro Gly Lys Ala Pro
    50                  55                  60

Lys Ser Leu Ile Tyr Ala Ala Ser Ser Leu Gln Ser Gly Val Pro Ser
65                  70                  75                  80

Lys Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser
                85                  90                  95

Ser Leu Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Tyr Asn
            100                 105                 110

Ser Phe Pro Leu Thr Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
        115                 120                 125
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<210> 26

<211> 469

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide

<400> 26

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gaagatctcc tgtaagggtt ctggatacag ctttaccagt tactggatcg gctgggtgcg    180

ccagatgccc gggaaaggcc tggagtggat ggggatcatc tatcctggtg actctgatac    240

cagatacagc ccgtccttcc aaggccaggt caccatctca gccgacaagt ccatcagcac    300

cgcctacctg cagtggagca gcctgaaggc ctcggacacc gccatgtatt actgtgcgag    360
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acggatggca gcagctggcc cctttgacta ctggggccag ggaaccctgg tcaccgtctc 420

ctcagcctcc accaagggcc catcggtctt ccccctggca ccctctagc 469

<210> 27

<211> 138

<212> PRT

<213> Homo sapiens

<400> 27

Met Gly Ser Thr Val Ile Leu Ala Leu Leu Leu Ala Val Leu Gln Gly
1 5 10 15

Val Cys Ala Glu Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
20 25 30

Pro Gly Glu Ser Leu Lys Ile Ser Cys Lys Gly Ser Gly Tyr Ser Phe
35 40 45

Thr Ser Tyr Trp Ile Gly Trp Val Arg Gln Met Pro Gly Lys Gly Leu
50 55 60

Glu Trp Met Gly Ile Ile Tyr Pro Gly Asp Ser Asp Thr Arg Tyr Ser
65 70 75 80

Pro Ser Phe Gln Gly Gln Val Thr Ile Ser Ala Asp Lys Ser Ile Ser
85 90 95

Thr Ala Tyr Leu Gln Trp Ser Ser Leu Lys Ala Ser Asp Thr Ala Met
100 105 110

Tyr Tyr Cys Ala Arg Arg Met Ala Ala Ala Gly Pro Phe Asp Tyr Trp
115 120 125

Gly Gln Gly Thr Leu Val Thr Val Ser Ser
130 135

<210> 28

<211> 466

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide

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taccactgga ggaatagtga tgacgcagtc tccagccacc ctgtctgtgt ctccagggga 120

aagagccacc ctctcctgca ggaccagtca gagtattggc tggaacttag cctggtacca 180

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cagttttggc caggggaccg agctggagat caaacgaact gtggctgcac catctgtctt 420

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<210> 29

<211> 128

<212> PRT

<213> Homo sapiens

<400> 29

Met Arg Val Pro Ala Gln Leu Leu Phe Leu Leu Leu Leu Trp Leu Pro
1 5 10 15

Asp Thr Thr Gly Gly Ile Val Met Thr Gln Ser Pro Ala Thr Leu Ser
20 25 30

Val Ser Pro Gly Glu Arg Ala Thr Leu Ser Cys Arg Thr Ser Gln Ser
35 40 45

Ile Gly Trp Asn Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro
50 55 60

Arg Leu Leu Ile Tyr Gly Ala Ser Ser Arg Thr Thr Gly Ile Pro Ala
65 70 75 80

Arg Phe Ser Gly Ser Gly Ser Gly Thr Glu Phe Thr Leu Thr Ile Ser
85 90 95

Ser Leu Gln Ser Glu Asp Ser Ala Val Tyr Tyr Cys Gln His Tyr Asp
100 105 110

Asn Trp Pro Met Cys Ser Phe Gly Gln Gly Thr Glu Leu Glu Ile Lys
115 120 125

<210> 30

<211> 487

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide

<400> 30

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ctctagc 487

<210> 31

<211> 144

<212> PRT

<213> Homo sapiens

<400> 31

Met Glu Phe Gly Leu Cys Trp Ile Phe Leu Val Ala Leu Leu Arg Gly
1 5 10 15

Val Gln Cys Gln Val Gln Leu Val Glu Ser Gly Gly Gly Val Val Gln
20 25 30

Pro Gly Arg Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe
35 40 45

Ile Ser Tyr Gly Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu
50 55 60

Glu Trp Val Ala Val Ile Ser Tyr Asp Gly Ser Asn Lys Tyr Tyr Ala
65 70 75 80

Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn
85 90 95

Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val
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Tyr Tyr Cys Ala Arg Val Leu Val Gly Ala Leu Tyr Tyr Tyr Asn Tyr
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Tyr Gly Met Asp Val Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser
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caaccggttc tctggagtgc cagataggtt cagtggcagc gggtcaggga cagatttcac 300

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<212> PRT

<213> Homo sapiens

<400> 33

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Val Thr Pro Gly Gln Pro Ala Ser Ile Ser Cys Lys Ser Ser Gln Ser
35 40 45

Leu Leu His Ser Asp Gly Lys Thr Phe Leu Tyr Trp Tyr Leu Gln Lys
50 55 60

Pro Gly Gln Pro Pro Gln Leu Leu Ile Tyr Glu Val Ser Asn Arg Phe
65 70 75 80

Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe
85 90 95

Thr Leu Lys Ile Ser Arg Val Glu Ala Glu Asp Val Gly Leu Tyr Tyr
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Cys Met Gln Ser Ile Gln Leu Pro Leu Thr Phe Gly Gly Gly Thr Lys
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Val Glu Ile Lys

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Arg Ser Pro Ala Leu Pro Phe Val Ser
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